OIPE

RAW SEQUENCE LISTING   DATE: 07/05/2001
PATENT APPLICATION: US/09/816,756A   TIME: 13:06:24

Input Set : **A:\SEQ LST - revised Missing Parts response.txt**
Output Set: **N:\CRF3\07032001\I816756A.raw**

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   3 <110> APPLICANT: University of Connecticut
   4        Mayer, Bruce
   6 <120> TITLE OF INVENTION: Coiled-Coil Mediated Heterodimerization Functional
Interaction Trap
   8 <130> FILE REFERENCE: 883933.0062
C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/816,756A
  11 <141> CURRENT FILING DATE: 2001-03-24
  13 <150> PRIOR APPLICATION NUMBER: 60/141,896
  14 <151> PRIOR FILING DATE: 1999-06-30
  16 <150> PRIOR APPLICATION NUMBER: PCT/US00/17929
  17 <151> PRIOR FILING DATE: 2000-06-29
  19 <160> NUMBER OF SEQ ID NOS: 2
  21 <170> SOFTWARE: PatentIn version 3.0
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ERRORED SEQUENCES

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  40 <210> SEQ ID NO: 2
  41 <211> LENGTH: 165
  42 <212> TYPE: DNA
  43 <213> ORGANISM: Artificial Sequence
  45 <220> FEATURE:
  47 <223> OTHER INFORMATION: Kozak translation start site, follwed by a Myc-epitope,
followed
  48        by a WIN-ZIP-B1 synthetic amphipathic helix, followed by an in-fr
  49        ame BamHI cloning site
  51 <400> SEQUENCE: 2
  52 accatggagc aaaagctcat ttctgaagag gacttgaatg aaggatctac catgtccgtt      60
  54 gacgaactgc aggctgaggt tgaccagctg caggacgaga attacgctct gaagaccaag     120
  56 gttgcgcagc tgcgtaaaaa ggtggaaaag ctgggaggag gatcc                     165
E--> 60 c&LdOc:.stmLibl:891220.1 06/12/01
```

VERIFICATION SUMMARY
PATENT APPLICATION: **US/09/816,756A**

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```
L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:60 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:184 SEQ:2
L:60 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:24
L:60 M:112 C: (48) String data converted to lower case,
L:60 M:252 E: No. of Seq. differs, <211>LENGTH:Input:165 Found:184 SEQ:2
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